

R. Swartz

1641

#6
MB
06/09/00

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/461,774

DATE: 05/19/2000
TIME: 18:58:25

Input Set: I461774.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

ENTERED

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JUN 08 2000

1 <110> APPLICANT: CHAN, Lily
2 CHUNG, Maxey Ching Ming
3 LIM, Renee Lay Hong
4 <120> TITLE OF INVENTION: Bacterial-derived molecules and therapeutic and
5 diagnostic uses therefor
6 <130> FILE REFERENCE: 1781-0180P
7 <140> CURRENT APPLICATION NUMBER: US/09/461,774
8 <141> CURRENT FILING DATE: 1999-12-15
9 <160> NUMBER OF SEQ ID NOS: 22
10 <170> SOFTWARE: PatentIn Ver. 2.1
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25 ggc atg gac aag ctg gcc gac acc gtg cgg gtg acg ctg ggg ccg cgc 96
26 Gly Met Asp Lys Leu Ala Asp Thr Val Arg Val Thr Leu Gly Pro Arg
27 20 25 30
28 ggc cgg cat gtg gtg ctg gcc aag gcg ttt ggc gga ccc acg gtt acc 144
29 Gly Arg His Val Val Leu Ala Lys Ala Phe Gly Gly Pro Thr Val Thr
30 35 40 45
31 aac gac ggc gtc acg gtg gca cgt gag atc gag ctg gaa gat ccg ttt 192
32 Asn Asp Gly Val Thr Val Ala Arg Glu Ile Glu Leu Glu Asp Pro Phe
33 50 55 60
34 gaa gac ttg ggc gcc cag ctg gtg aag tcg gtg gcc acc aag acc aac 240
35 Glu Asp Leu Gly Ala Gln Leu Val Lys Ser Val Ala Thr Lys Thr Asn
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37 gat gtt gcc ggt gac ggc acc acc acc gca acc atc ttg gcg cag gca 288
38 Asp Val Ala Gly Asp Gly Thr Thr Thr Ala Thr Ile Leu Ala Gln Ala
39 85 90 95
40 ctg atc aag ggc ggc ctg agg cta gtg gcc gcc ggc gtc aac ccg gtc 336
41 Leu Ile Lys Gly Gly Leu Arg Leu Val Ala Ala Gly Val Asn Pro Val
42 100 105 110
43 gcg ctc ggc gtg gga atc ggc aag gcc gcc gac gcg gta ttc gag gcg 384
44 Ala Leu Gly Val Gly Ile Gly Lys Ala Ala Asp Ala Val Phe Glu Ala

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49	gtg	gcg	acg	gtg	tcc	tcg	cgc	gac
50	Val	Ala	Thr	Val	Ser	Ser	Arg	Asp
51		145			150			155
52	gaa	gcg	atg	aac	aag	gtc	ggc	cac
53	Glu	Ala	Met	Asn	Lys	Val	Gly	His
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55	tcc	tcc	acg	ctg	ggc	acc	gag	ttg
56	Ser	Ser	Thr	Leu	Gly	Thr	Glu	Leu
57			180				185	
58	cac	aaa	ggc	ttc	ttg	tcg	gca	tac
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61	cag	gcg	gtg	ctc	gag	gac	gcg	ttg
62	Gln	Ala	Val	Leu	Glu	Asp	Ala	Leu
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64	agc	tcg	ctt	ccc	gat	ctg	ttg	cca
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66		225			230			235
67	ggg	aag	cca	cta	ctg	atc	gtg	gct
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69			245				250	
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71	Ala	Thr	Leu	Val	Val	Asn	Ala	Ile
72			260				265	
73	gtc	aag	ggg	ccg	tac	ttc	ggg	gac
74	Val	Lys	Gly	Pro	Tyr	Phe	Gly	Asp
75		275			280			285
76	ctg	gcg	gtg	gtg	acg	ggg	ggc	cac
77	Leu	Ala	Val	Val	Thr	Gly	Gly	His
78		290			295			300
79	gtg	ctg	cgc	gag	gtg	ggc	ttg	gag
80	Val	Leu	Arg	Glu	Val	Gly	Leu	Glu
81		305			310			315
82	gtg	gtc	agc	aag	gac	gac	acg	gtc
83	Val	Val	Ser	Lys	Asp	Asp	Thr	Val
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85	gaa	gcg	gtg	gcc	aac	cgg	gcg	aac
86	Glu	Ala	Val	Ala	Asn	Arg	Ala	Asn
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88	agc	gat	tcg	gat	tgg	gat	cgg	gaa
89	Ser	Asp	Ser	Asp	Trp	Asp	Arg	Glu
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91	ctg	gcc	ggc	ggg	gtt	gct	gtc	atc
92	Leu	Ala	Gly	Gly	Val	Ala	Val	Ile
93		370			375			380
94	gca	ctc	aag	gag	cgc	aag	gaa	agc

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98   Lys Ala Ala Val Glu Glu Gly Ile Val Pro Gly Gly Gly Ala Ser Leu
99                               405                               410                               415
100  atc cac cag gcc cgc aag gcg ctg acc gaa ctg cgt gcg tcg ctg acc 1296
101  Ile His Gln Ala Arg Lys Ala Leu Thr Glu Leu Arg Ala Ser Leu Thr
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103  ggt gac gag gtc ctc ggt gtc gac gtg ttc tcc gaa gcc ctt gcc gcg 1344
104  Gly Asp Glu Val Leu Gly Val Asp Val Phe Ser Glu Ala Leu Ala Ala
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106  ccg ttg ttc tgg atc gcc gcc aac gct ggc ttg gac ggc tcg gtg gtg 1392
107  Pro Leu Phe Trp Ile Ala Ala Asn Ala Gly Leu Asp Gly Ser Val Val
108                               450                               455                               460
109  gtc aac aag gtc agc gag cta ccc gcc ggg cat ggg ctg aac gtg aac 1440
110  Val Asn Lys Val Ser Glu Leu Pro Ala Gly His Gly Leu Asn Val Asn
111  465                               470                               475                               480
112  acc ctg agc tat ggt gac ttg gcc gct gac ggc gtc atc gac ccg gtc 1488
113  Thr Leu Ser Tyr Gly Asp Leu Ala Ala Asp Gly Val Ile Asp Pro Val
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115  aag gtg act agg tcg gcg gtg ttg aac gcg tca tcg gtt gcc cgg atg 1536
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118  gta ctc acc acc gag acg gtc gtg gtc gac aag ccg gcc aag gca gaa 1584
119  Val Leu Thr Thr Glu Thr Val Val Val Asp Lys Pro Ala Lys Ala Glu
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126  <212> TYPE: PRT
127  <213> ORGANISM: Mycobacterium tuberculosis
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135  Asn Asp Gly Val Thr Val Ala Arg Glu Ile Glu Leu Glu Asp Pro Phe
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137  Glu Asp Leu Gly Ala Gln Leu Val Lys Ser Val Ala Thr Lys Thr Asn
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141  Leu Ile Lys Gly Gly Leu Arg Leu Val Ala Ala Gly Val Asn Pro Val
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RAW SEQUENCE LISTING
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148	145					150					155				160	
149	Glu	Ala	Met	Asn	Lys	Val	Gly	His	Asp	Ser	Val	Val	Ser	Val	Lys	Glu
150					165					170					175	
151	Ser	Ser	Thr	Leu	Gly	Thr	Glu	Leu	Glu	Phe	Thr	Glu	Gly	Ile	Gly	Phe
152				180						185				190		
153	His	Lys	Gly	Phe	Leu	Ser	Ala	Tyr	Phe	Val	Thr	Asp	Phe	Asp	Asn	Gln
154			195					200					205			
155	Gln	Ala	Val	Leu	Glu	Asp	Ala	Leu	Ile	Leu	Leu	His	Gln	Asp	Lys	Ile
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189	Thr	Leu	Ser	Tyr	Gly	Asp	Leu	Ala	Ala	Asp	Gly	Val	Ile	Asp	Pro	Val
190				485						490					495	
191	Lys	Val	Thr	Arg	Ser	Ala	Val	Leu	Asn	Ala	Ser	Ser	Val	Ala	Arg	Met
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194				515					520					525		

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211      ggg ttg ctg ctc ggc gga ctg ctc cta gca gcg ccc ccg ttg gcc ggc      96
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221      Ser Glu Ile Arg Thr Ala Arg Cys Ala Thr Val Ser Val Pro Val Asp
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223      tac gac caa ccc ggt ggg aca caa gcg aag ttg gcg gtg atc cgc gtc      288
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VERIFICATION SUMMARY
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DATE: 05/19/2000

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Input Set: I461774.RAW

Line ? Error/Warning

Original Text

306 W Invalid/Missing Amino Acid Numbering

515

520